

Specificity Table / Spezifitätstabelle

Mix	5'-Primer [®]	3'-Primer [®]	bp	Specificity / Spezifität
1	88 GAC	169 AAC	120	DQB1*05:01:01:01-01:19,07,11:01-12,18,20-22,25,27,29-32,44,45,48,49,51,54,55,61-63,68-76,80-82,84,88,89
2	88 GAC	169 GCT	120	DQB1*05:02:01:02:11, 05:01,05:02,14,17,19,26,33-37,46,47,53,57,59,65,79,83,86,87Q
3	88 GAC	170 CGT 170 CAT	125	DQB1*05:03:01:01-03:16,06:01,06:02,08-10,13,15,16,23,24,28,38-41N,42,43,50,56,58,60,64,66,67,78,85 / *06:162
4	40 GCC	198 ATG	200	DQB1*05:04,52,77
5	404 TGA	573 GAT	800 [®] KB (410bp)	DQB1*02:01:01-01:19,04,05?,07:01?-09?,13?-18N?,19?,20N?-52?,53
6	63 ACA	170 CGG	150	DQB1*02:01:01-01:06,01:08-01:19,02:01,02:02,04-14:02w,15,16,18N-20N,21-33w,34-41,42-53Q
7	404 TGG	503 CTC	140	DQB1*02:02:01,02:02,03,05?,06w,10-12,13?-18N?,19?,20N?-26?,28?-47?,49?-52?
8	63 ACA	170 CGT	150	DQB1*02:03 / *04:02:02
9	406 ACG	573 GAT	795 [®] KB (410bp)	DQB1*02:04, 05?
10	38 GGC	198 ACT	200	DQB1*03:01:01:01-01:28,04:01,04:02,09-10:02,12,14:01,14:02,16,19,21,22,24,27-29,35,36,42,44,46-60,70,71,73-78,80,82-84N,92-94,101-103,108,109,114-116,118N-122,127-131,133-135,139,140,143,144,147,148,150-152,154,157-160,162-167 / *06:58,122,148
11	171 GAC	230 CCG	100	DQB1*03:01:01:01-01:13w,01:14-01:27,03:02:01-03:08w,03:09-03:13,09-10:02,12,13,15-17:02,19-22,24,26-31,33-36,38-44,46-60,65,69,71-79,82-84N,86-90N,91Q-95N,96-99Q,100-105,108,109,111-118N,119-124,126-129w,130w,131,133-137,139-145,147-152,154-160,162-167 / *06:66,96
12	134 GGA	198 ACT	105	DQB1*03:01:01:01-01:28,04:01,04:02,09,16,19,21,22,24,27-29,35,36,42,44,46-60,69,71,73-78,80,82-84N,92-94,101-103,108,109,113-116,118N-122,127-131,133-135,139,140,142-144,147,148,150-152,154,157-160,162-167 / *06:145
13	114 GCA	208 CCT	135	DQB1*03:01:01:01-01:03,01:03-01:06,01:08-01:28,02:01,02:02, 02:04-02:17,03:02:01-03:13,04:01,04:02,05:03,05:04,07,09-19,21,22, 24, 26-58w,59,60,62-64,66N-71,74,76-84N,85-90N,91Q-95N,96-99Q,101-107w,108-111,113-118Nw,119-131,133-136,138-155,157-161,163-167 / *06:04:08
14	114 GCA	170 CGG	95	DQB1*03:02:01,02:02,02:04-02:09,02:11-02:14w,02:15-02:17,04:01,04:02,05:03,05:04,07,08,11,14:01,14:02,18, 32,37,45,62-64,66N-68,70,80,81,85,106,107,125,138,146,153,161
15	146 GGC	250 CTG	145	DQB1*03:02:01-02:11w,02:12-02:17,03:02:01-03:13,04:02,05:01-05:04, 06,08,10:01,10:02w-12, 14:02,15,17:01-18,20,25,26,30-34,37-41,43,62-64w,65,66N-68, 70,72,79,81,85-89,91Q,95N-99Q,100,104-107,110-112,117,123-126,132,136-138,141,145,146,149,153,155,156,161 / *04:01:01-02:05w,02:06,02:07-08,10,12-25N,26,27
16	77 TCT	170 CGT	135	DQB1*03:03:02:01-03:13,06,12,15,20,25,26,30,31,33,34,38-41,43,65, 74,79,86-90N,91Q,95N-99Q,104,105,111-113,117,123,124,126,136,137,141,145,149,155,156 / *02:03 / *04:03:01,03:02 / *06:03:10,51:01,66,96
17	77 TTA	170 CGG	135	DQB1*03:04:01,04:02,14:01,14:02,80,138
18	75 CGG	170 CGG	135	DQB1*03:05:01-05:04,61,132
19	77 TCT	231 ACG	195	DQB1*03:06, 25 / *04:03:01, 03:02
20	113 CGC	146 CCA 208 CCC	75 135	DQB1*03:07 DQB1*03:08,137 / *06:02:01-02:07,02:09-02:23,03:01-03:18,08:01-08:03,10-14:02,16, 19:01-21,23,26N,29,31,33,40,41,44,46,47,49,50,59-62w,63-65,67,68,70-75N,76,77N,78,80-84,87,90-92,95-97,106,107,109-112N,113-117,119,122-128,130,131,133,134,136-138,141,143,145-148,150-152,154,156,159,161-163,165,166
21	134 GGA	503 CGG	650	DQB1*03:09 (sometimes a weak unspecific band at about 70bp)
22	40 GCC 79 TTC 112 ACA	198 ACT	200 160 125	DQB1*03:11,26 / *05:01:01:01-01:18w,01:19,02:01-02:11,03:01:01-03:16,05:01-41N,42-50,53-76,78-87Q,88,89 / *06:05:01,05:02,06,20,31,156 DQB1*03:18 DQB1*03:20 / *05:28
23	38 GGC	77 CAA	80	DQB1*03:12,70,74,78 / *06:122
24	114 GCA	167 CAA 189 TGC 200 GGT	95 115 125	DQB1*03:25,130w / *04:01:03 DQB1*03:15 DQB1*03:13 (sometimes a weak unspecific band at about 900 bp)
25	171 GAC	389 ACT 397 CCA 508 GAT	500 505 620	DQB1*03:22 DQB1*03:21 DQB1*03:24,79
26	68 GCT	231 ACG	205	DQB1*04:01:01-01:04,05-08,14-17

Mix	5'-Primer [Ⓞ]	3'-Primer [Ⓞ]	bp	Specificity / Spezifität
27	68 GCG	231 ACG	205	DQB1*04:02:01-02:07,03:01,04,09-13,18-20w,21-25N,26,27 / *03:132
28	141 TAT	216 GCT	115	DQB1*06:01:01-01:11,01:14,45,53,54N-57,98,99:02-102N,103-105,108,120,132,140,153,157,167
29	88 GAT	144 GCG	95	DQB1*06:02:01-02:04,02:06-02:15,02:17,02:18,02:20-02:23,09:02,10,11:01-11:03,13:01,15:01,15:02,16,19:02,20,22:03,24,29,33,37,46-51:02,68,70-75N,76,77N-84,94,95w-97,99:01,106,107,109,111,112N-118:01,122,124-127,130,131,136-138,146,147,150,151,159,161,166 / *03:02:05,23
30	88 GAT	170 CAT	125	DQB1*06:01:08,02:01-02:11w,02:12-02:23,11:01,11:03,15:01,15:02,19:01-20,24,33,37,46-50,51:02,68,70-75N,76,77N-84,95,97,106w,107,109,111,112N,113-118:02,122,124-127,136-138,146,147,150-152,159,161,163,166 / *03:23
31	171 GAT	259 GAA	130	DQB1*06:01:08,02:01-02:21w,02:22-03:09,03:11-03:17w,03:18,11:01-11:03,14:01,14:02,16,19:01-20,23,24,26N,28,30-33,37,40,41,44,47-50,51:02,59-62,64,65,67,68,70-72w,73w-75N,76w,77N-84,87w,90,91,95,97,106,107,109-112N,113-117,122,124-128,131,134,136-138,141,143,144N,145,147,148,150-152,154,156,159,161-163w,165,166
32	81 GTA	208 CCT	170	DQB1*06:04:01-04:10,05:01,05:02,07:01,07:02,09:01-09:03,09:05,09:06,18:01,18:02,22:02,25,27:01,27:02,32,34,36,38,39,52,58,66,69,85,86,88,89,93,94,118:01,118:02,121,129,135,142,155,158N,160,164
33	75 CGT	170 CAA	135	DQB1*06:04:01-04:10,05:01,06,08:01-09:06,12-13:02,17,18:01,18:02,21-22:03,27:01,27:02,34,36,38,39,42,52,55,58,69,85,86,88,89,93,94w,119,121,129,135,142,149,155,158N,160,164
34	88 GAC	257 ACC	210	DQB1*06:04:01-04:05,04:07-04:10,07:01,07:02,17,21,25,34,36,38,39,52,58,69,85,86,89,92,93,135,149,155,158N,160,164
35	77 TCT	257 ACC	220	DQB1*06:04:01-04:05,04:07-05:02,06?,07:01,07:02,09:01-09:06,12,15:01,15:02,17,21-22:03,25,34,36,38,39,42,46,52,58,66,69,85,86,88,89,92-94,118:01-119,121,135,142,149,155,158N,160,164
36	40 GCA	257 ACC	260	DQB1*06:04:01-04:05,04:07-04:10,07:01,07:02,09:01-09:06,12,15:01,15:02,17,21-22:03,25,34,36,38,39,42,46,52,66,69,86,88,89,92-94,118:01-119,121,135,142,149,155,158N,160,164
37	26 GTT	211 GGT	225	DQB1*06:02:01-02:23,10,13:01-16,19:01-20,22:01-23,29,33,37,46-51:01w,51:02,68-75N,76,77N-84,95-97,106,107,109,111,112N-117,119,122-127,130,131w,136-139,146,147,150-152,156,159,161-163,166 / *03:30,72,100
38	26 GTA	259 GAA	275	DQB1*06:03:01-03:17w,03:18,06?,08:01-08:03,11:01-11:03,18:01,18:02,26N,27:01-28,30-32,40,41,44,45,59-65,67,87,90,91,110,128,133,134,141,143,144N,145,148,154,165 / *05:73
39	114 GCG	169 GCT	95	<i>DQB1*05:05:01 / *06:10,25,130</i>
40	26 GTA	88 GTA	105	DQB1*06:05:01,05:02,06?,09:01-09:06,11:01-11:03,12,18:01,18:02,42,45,66,88,94,118:01,118:02,121,129,142 / *03:01:01:01-29,31-37,39-46w,47-51,53-66N,67-71,73-84N,85-90N,91Q-95N,96-98,101-118N,119-126,128-131,133-158w,159,160w-167 / *04:04,05 / 05:04,52,77
41	26 GTT	88 GTG	105	<i>DQB1*06:14:01,14:02,69,156,162 / *05:38,62</i>
42	26 GTT	178 GTT	195	<i>DQB1*06:16</i>
43	77 TCT	211 GGC	175	<i>DQB1*06:17,24,30,42</i>
44	26 GTT	141 CGA	155	DQB1*06:02:05,02:14w,19:01,139 / *03:30,72,100,132 / *04:01:01-02:04,02:06,02:08-03:02,06-08,10,11w-25N,26,27 / *05:38
45	40 GCC 85 CCT	144 GCG	145 100	<i>DQB1*06:20,31</i> <i>DQB1*06:26N,156 / *05:03:12</i>
46	77 GGG 33 TTA 91 ACG	144 GCG	110 150 95	<i>DQB1*06:23,156,162 / *05:03:12</i> <i>DQB1*06:33</i> <i>DQB1*06:40</i>
47	554 CAT	573 GAC	650	DQB1*03:02:01-03:13,05:01-05:04,06?-08?,11?-15?,17:01?-18?,19,20?,23?,25?,26?,30?,31-34?,37?-41?,43?,45?,48?,52?-66N?,67?-72?,74?-79?,81?,82?,85?-90N?,91Q,95N-99Q,100,101?-112?,117,118N?,120?-138?,140?-163?,165?-167? / *04:01:01,01:02?-01:04?,02:01,02:02?-02:04?,02:05,02:06?-02:08?,03:01?-25N?,26?,27?
48				Kontaminationskontrolle / Contamination control

Ⓞ The codon at the 3'-end of the primer. The number given is the nucleotide position.
Das Codon am 3'-Ende des Primers. Die Nummer gibt die Nucleotidposition an.

Ⓠ The control band is smaller than the specific band (410 bp) / Die Kontrollbande ist kleiner als die spezifische Bande (410 bp)

. Alleles in italics: Positive reactions could not be verified experimentally. The correct reactivity of the primers was tested in other mixes if possible. /
Kursiv gedruckte Allele: Die positiven Reaktionen konnten nicht experimentell überprüft werden. Die einzelnen Primer wurden aber soweit möglich in anderen Mixen auf korrekte Reaktivität überprüft.

w weak reaction / schwache Reaktion

? no sequence information available for primer binding site / keine Sequenzinformation für Primerbindungsstelle vorhanden

The following alleles cannot be detected / Die folgenden Allele werden nicht erkannt: / : DQB1*06:01:12,35,43

